CRF Errors Corrected by the STIC Systems Branch

Serial Number: 10/030,846 ENTERED

CRF Processing Date: 2/12/2003

Edited by:

Verified by: (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☑ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____.
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

PCT10

RAW SEQUENCE LISTING DATE: 02/12/2003
PATENT APPLICATION: **US/10/030,846** TIME: 08:03:51

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\02122003\J030846.raw**

<110> APPLICANT: Larose, Anne-Marie
Leblanc, Benoet
Camato, Rino
<120> TITLE OF INVENTION: METHOD FOR NORMALIZING THE RELATIVE INTENSITIES OF DETECTION SIGNALS IN HYBRIDIZATION ARRAYS
<130> FILE REFERENCE: 9555.127USWO
<140> CURRENT APPLICATION NUMBER: US 10/030,846
<141> CURRENT FILING DATE: 2002-01-11
<150> PRIOR APPLICATION NUMBER: PCT/CA01/01860
<151> PRIOR FILING DATE: 2001-12-21
<150> PRIOR APPLICATION NUMBER: CA 2,327,527
<151> PRIOR FILING DATE: 2000-12-27
<160> NUMBER OF SEQ ID NOS: 3
<170> SOFTWARE: PatentIn Ver. 2.1
<210> SEQ ID NO: 1
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Oligonucleotide
<400> SEQUENCE: 1
cttatgaccc gcacttactc g 21
<210> SEQ ID NO: 2
<211> LENGTH: 40
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Oligonucleotide
<400> SEQUENCE: 2
cccgagccgc ctggataccg cagctaggaa taatggaata 40
<210> SEQ ID NO: 3
<211> LENGTH: 40
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Oligonucleotide
<400> SEQUENCE: 3
tctcgattcc gtgggtggtg gtgcatggcc gttcttagtt 40

VERIFICATION SUMMARY DATE: 02/12/2003
PATENT APPLICATION: **US/10/030,846** TIME: 08:03:52

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\02122003\J030846.raw**